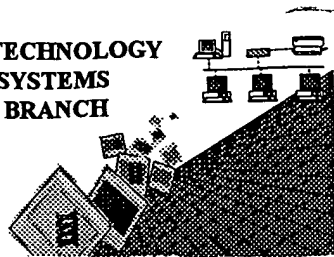# RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/018,383_
Source: _Puzllo_
Date Processed by STIC: _6/4/2002_

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
 PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
 PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002

PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/018,383

DATE: 06/04/2002
TIME: 14:56:06

Input Set : A:\X11600.ST25.txt
Output Set: N:\CRF3\06042002\J018383.raw

Does Not Comply
Corrected Diskette Needed

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   3 <110> APPLICANT: Becker, Gerald
   4        Hale, John
   5        Heath, William
   6        Johnstone, Edward
   7        Little, Sheila
   8        Tu, Yuan
   9        Yeh, Wu-Kuang
  10        Yin, Tinggui
  12 <120> TITLE OF INVENTION: Amyloid Precursor Protein Protease and Related Nucleic Acid
Compounds
  14 <130> FILE REFERENCE: X-11600
C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/018,383
C--> 16 <141> CURRENT FILING DATE: 2002-03-05
  16 <160> NUMBER OF SEQ ID NOS: 5
  18 <170> SOFTWARE: PatentIn version 3.1
  20 <210> SEQ ID NO: 1
  21 <211> LENGTH: 1683
  22 <212> TYPE: DNA
  23 <213> ORGANISM: Homo sapiens
  25 <220> FEATURE:
  26 <221> NAME/KEY: CDS
  27 <222> LOCATION: (25)..(1629)
  28 <223> OTHER INFORMATION:
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  32 <221> NAME/KEY: misc_feature
  33 <222> LOCATION: (25)..(1629)
  34 <223> OTHER INFORMATION: n = any nucleotide A, C, G, or T
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  39                             Met Leu Arg Arg Arg Gly Ser Pro Gly
  40                              1               5
  42 atg ggt gtg cat gtg ggt gca gcc ctg gga gca ctg tgg ttc tgc ctc        99
  43 Met Gly Val His Val Gly Ala Ala Leu Gly Ala Leu Trp Phe Cys Leu
  44 10               15                  20                  25
  46 aca gga gcc ctg gag gtc cag gtc cct gaa gac cca gtg gtg gca ctg       147
  47 Thr Gly Ala Leu Glu Val Gln Val Pro Glu Asp Pro Val Val Ala Leu
  48               30                  35                  40
  50 gtg ggc acc gat gcc acc ctg tgc tgc tcc ttc tcc cct gag cct ggc       195
  51 Val Gly Thr Asp Ala Thr Leu Cys Cys Ser Phe Ser Pro Glu Pro Gly
  52               45                  50                  55
  54 ttc agc ctg gca cag ctc aac ctc atc tgg cag ctg aca gat acc aaa       243
  55 Phe Ser Leu Ala Gln Leu Asn Leu Ile Trp Gln Leu Thr Asp Thr Lys
  56               60                  65                  70
  58 cag ctg gtg cac agc ttt gct gag ggc cag gac cag ggc agc gcc tat       291
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RAW SEQUENCE LISTING   DATE: 06/04/2002
PATENT APPLICATION: US/10/018,383   TIME: 14:56:06

Input Set : A:\X11600.ST25.txt
Output Set: N:\CRF3\06042002\J018383.raw

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59 Gln Leu Val His Ser Phe Ala Glu Gly Gln Asp Gln Gly Ser Ala Tyr
60      75              80              85
62 gcc aac cgc acg gcc ctc ttc ccg gac ctg ctg gca cag ggc aac gca      339
63 Ala Asn Arg Thr Ala Leu Phe Pro Asp Leu Leu Ala Gln Gly Asn Ala
64 90              95              100             105
66 tcc ctg agg ctg cag cgc gtg cgt gtg gcg gac gag ggc agc ttc acc      387
67 Ser Leu Arg Leu Gln Arg Val Arg Val Ala Asp Glu Gly Ser Phe Thr
68              110             115             120
70 tgc ttc gtg agc atc cgg gat ttc ggc agc gct gcc gtc agc ctg cag      435
71 Cys Phe Val Ser Ile Arg Asp Phe Gly Ser Ala Ala Val Ser Leu Gln
72              125             130             135
74 gtg gcc gct ccc tac tcg aag ccc agc atg acc ctg gag ccc aac aag      483
75 Val Ala Ala Pro Tyr Ser Lys Pro Ser Met Thr Leu Glu Pro Asn Lys
76              140             145             150
78 gac ctg cgg cca ggg gac acg gtg acc atc acg tgc tcc agc tac cag      531
79 Asp Leu Arg Pro Gly Asp Thr Val Thr Ile Thr Cys Ser Ser Tyr Gln
80      155             160             165
82 ggc tac cct gag gct gag gtg ttc tgg cag gat ggg cag ggt gtg ccc      579
83 Gly Tyr Pro Glu Ala Glu Val Phe Trp Gln Asp Gly Gln Gly Val Pro
84 170             175             180             185
86 ctg act ggc aac gtg acc acg tcg cag atg gcc aac gag cag ggc ttg      627
87 Leu Thr Gly Asn Val Thr Thr Ser Gln Met Ala Asn Glu Gln Gly Leu
88              190             195             200
90 ttt gat gtg cac agc atc ctg cgg gtg gtg ctg ggt gca aat ggc acc      675
91 Phe Asp Val His Ser Ile Leu Arg Val Val Leu Gly Ala Asn Gly Thr
92              205             210             215
94 tac agc tgc ctg gtg cgc aac ccc gtg ctg cag cag gat gcg cac agc      723
95 Tyr Ser Cys Leu Val Arg Asn Pro Val Leu Gln Gln Asp Ala His Ser
96              220             225             230
98 tct gtc acc atc aca ccc cag aga agc ccc aca gga gcc gtg gag gtc      771
99 Ser Val Thr Ile Thr Pro Gln Arg Ser Pro Thr Gly Ala Val Glu Val
100     235             240             245
102 cag gtc cct gag gac ccg gtg gtg gcc cta gtg ggc acc gat gcc acc     819
103 Gln Val Pro Glu Asp Pro Val Val Ala Leu Val Gly Thr Asp Ala Thr
104 250             255             260             265
106 ctg cgc tgc tcc ttc tcc ccc gag cct ggc ttc agc ctg gca cag ctc     867
107 Leu Arg Cys Ser Phe Ser Pro Glu Pro Gly Phe Ser Leu Ala Gln Leu
108             270             275             280
110 aac ctc atc tgg cag ctg aca gac acc aaa cag ctg gtg cac agt ttc     915
111 Asn Leu Ile Trp Gln Leu Thr Asp Thr Lys Gln Leu Val His Ser Phe
112             285             290             295
114 acc gaa ggc cgg gac cag ggc agc gcc tat gcc aac cgc acg gcc ctc     963
115 Thr Glu Gly Arg Asp Gln Gly Ser Ala Tyr Ala Asn Arg Thr Ala Leu
116             300             305             310
118 ttc ccg gac ctg ctg gca caa ggc aat gca tcc ctg agg ctg cag cgc     1011
119 Phe Pro Asp Leu Leu Ala Gln Gly Asn Ala Ser Leu Arg Leu Gln Arg
120     315             320             325
122 gtg cgt gtg gcg gac gag ggc agc ttc acc tgc ttc gtg agc atc cgg     1059
123 Val Arg Val Ala Asp Glu Gly Ser Phe Thr Cys Phe Val Ser Ile Arg
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                    RAW SEQUENCE LISTING          DATE: 06/04/2002
                    PATENT APPLICATION:  US/10/018,383   TIME: 14:56:06

                    Input Set : A:\X11600.ST25.txt
                    Output Set: N:\CRF3\06042002\J018383.raw


       124 330                   335              340                  345
       126 gat ttc ggc agc gct gcc gtc agc ctg cag gtg gcc gct ccc tac tcg      1107
       127 Asp Phe Gly Ser Ala Ala Val Ser Leu Gln Val Ala Ala Pro Tyr Ser
       128           350              355                  360
       130 aag ccc agc atg acc ctg gag ccc aac aag gac ctg cgg cca ggg gac      1155
       131 Lys Pro Ser Met Thr Leu Glu Pro Asn Lys Asp Leu Arg Pro Gly Asp
       132           365              370                  375
       134 acg gtg acc atc acg tgc tcc agc tac cgg ggc tac cct gag gct gag      1203
       135 Thr Val Thr Ile Thr Cys Ser Ser Tyr Arg Gly Tyr Pro Glu Ala Glu
       136           380              385                  390
       138 gtg ttc tgg cag gat ggg cag ggt gtg ccc ctg act ggc aac gtg acc      1251
       139 Val Phe Trp Gln Asp Gly Gln Gly Val Pro Leu Thr Gly Asn Val Thr
       140           395              400                  405
W-->   142 acg tcg cag atg gcc aac gag cag ggc ttg ttt nnn nnn nnn nnn nnn      1299
W-->   143 Thr Ser Gln Met Ala Asn Glu Gln Gly Leu Phe Xaa Xaa Xaa Xaa Xaa
       144 410              415              420                  425
W-->   146 nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn nnn      1347
W-->   147 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
       148           430              435                  440
W-->   150 nnn nnn nnn nnn nnn nnn gat gcg cac ggc tct gtc acc atc aca ggg      1395
W-->   151 Xaa Xaa Xaa Xaa Xaa Xaa Asp Ala His Gly Ser Val Thr Ile Thr Gly
       152           445              450                  455
       154 cag cct atg aca ttc ccc cca gag gcc ctg tgg gtg acc gtg ggg ctg      1443
       155 Gln Pro Met Thr Phe Pro Pro Glu Ala Leu Trp Val Thr Val Gly Leu
       156           460              465                  470
       158 tct gtc tgt ctc att gca ctg ctg gtg gcc ctg gct ttc gtg tgc tgg      1491
       159 Ser Val Cys Leu Ile Ala Leu Leu Val Ala Leu Ala Phe Val Cys Trp
       160 475              480              485
       162 aga aag atc aaa cag agc tgt gag gag gag aat gca gga gct gag gac      1539
       163 Arg Lys Ile Lys Gln Ser Cys Glu Glu Glu Asn Ala Gly Ala Glu Asp
       164 490              495              500                  505
       166 cag gat ggg gag gga gaa ggc tcc aag aca gcc ctg cag cct ctg aaa      1587
       167 Gln Asp Gly Glu Gly Glu Gly Ser Lys Thr Ala Leu Gln Pro Leu Lys
       168           510              515                  520
       170 cac tct gac agc aaa gaa gat gat gga caa gaa ata gcc tga            1629
       171 His Ser Asp Ser Lys Glu Asp Asp Gly Gln Glu Ile Ala
       172           525              530
       174 ccatgaggac cagggagctg ctaccctcc ctacagctcc taccctctgg ctgc        1683
       177 <210> SEQ ID NO: 2
       178 <211> LENGTH: 534
       179 <212> TYPE: PRT
       180 <213> ORGANISM: Homo sapiens
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       183 <221> NAME/KEY: misc_feature
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       185 <223> OTHER INFORMATION: The 'Xaa' at location 421 stands for Lys, Asn, Arg, Ser,
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       186         , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
       187         Tyr, Trp, Cys, or Phe.
       189 <220> FEATURE:
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(handwritten) Xaa can only represent an actual amino acid, not a stop codon   6/4/02

RAW SEQUENCE LISTING   DATE: 06/04/2002
PATENT APPLICATION: US/10/018,383   TIME: 14:56:06

Input Set : **A:\X11600.ST25.txt**
Output Set: **N:\CRF3\06042002\J018383.raw**

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    192 <223> OTHER INFORMATION: The 'Xaa' at location 422 stands for Lys, Asn, Arg, Ser,
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    194       Tyr, Trp, Cys, or Phe.
    196 <220> FEATURE:
    197 <221> NAME/KEY: misc_feature
    198 <222> LOCATION: (423)..(423)
    199 <223> OTHER INFORMATION: The 'Xaa' at location 423 stands for Lys, Asn, Arg, Ser,
Thr, Ile
    200       , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
    201       Tyr, Trp, Cys, or Phe.
    203 <220> FEATURE:
    204 <221> NAME/KEY: misc_feature
    205 <222> LOCATION: (424)..(424)
    206 <223> OTHER INFORMATION: The 'Xaa' at location 424 stands for Lys, Asn, Arg, Ser,
Thr, Ile
    207       , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
    208       Tyr, Trp, Cys, or Phe.
    210 <220> FEATURE:
    211 <221> NAME/KEY: misc_feature
    212 <222> LOCATION: (425)..(425)
    213 <223> OTHER INFORMATION: The 'Xaa' at location 425 stands for Lys, Asn, Arg, Ser,
Thr, Ile
    214       , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
    215       Tyr, Trp, Cys, or Phe.
    217 <220> FEATURE:
    218 <221> NAME/KEY: misc_feature
    219 <222> LOCATION: (426)..(426)
    220 <223> OTHER INFORMATION: The 'Xaa' at location 426 stands for Lys, Asn, Arg, Ser,
Thr, Ile
    221       , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
    222       Tyr, Trp, Cys, or Phe.
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    226 <222> LOCATION: (427)..(427)
    227 <223> OTHER INFORMATION: The 'Xaa' at location 427 stands for Lys, Asn, Arg, Ser,
Thr, Ile
    228       , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
    229       Tyr, Trp, Cys, or Phe.
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    233 <222> LOCATION: (428)..(428)
    234 <223> OTHER INFORMATION: The 'Xaa' at location 428 stands for Lys, Asn, Arg, Ser,
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    236       Tyr, Trp, Cys, or Phe.
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    240 <222> LOCATION: (429)..(429)
    241 <223> OTHER INFORMATION: The 'Xaa' at location 429 stands for Lys, Asn, Arg, Ser,
Thr, Ile
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same eva

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242         ·, Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, (a stop codon,
243         Tyr, Trp, Cys, or Phe.
245 <220> FEATURE:
246 <221> NAME/KEY: misc_feature
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RAW SEQUENCE LISTING   DATE: 06/04/2002
PATENT APPLICATION: US/10/018,383   TIME: 14:56:06

          Input Set : A:\X11600.ST25.txt
          Output Set: N:\CRF3\06042002\J018383.raw

     247 <222> LOCATION: (430)..(430)
     248 <223> OTHER INFORMATION: The 'Xaa' at location 430 stands for Lys, Asn, Arg, Ser,
Thr, Ile
     249   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
     250   Tyr, Trp, Cys, or Phe.
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     254 <222> LOCATION: (431)..(431)
     255 <223> OTHER INFORMATION: The 'Xaa' at location 431 stands for Lys, Asn, Arg, Ser,
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     256   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
     257   Tyr, Trp, Cys, or Phe.
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     261 <222> LOCATION: (432)..(432)
     262 <223> OTHER INFORMATION: The 'Xaa' at location 432 stands for Lys, Asn, Arg, Ser,
Thr, Ile
     263   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
     264   Tyr, Trp, Cys, or Phe.
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     268 <222> LOCATION: (433)..(433)
     269 <223> OTHER INFORMATION: The 'Xaa' at location 433 stands for Lys, Asn, Arg, Ser,
Thr, Ile
     270   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
     271   Tyr, Trp, Cys, or Phe.
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     274 <221> NAME/KEY: misc_feature
     275 <222> LOCATION: (434)..(434)
     276 <223> OTHER INFORMATION: The 'Xaa' at location 434 stands for Lys, Asn, Arg, Ser,
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     277   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
     278   Tyr, Trp, Cys, or Phe.
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     282 <222> LOCATION: (435)..(435)
     283 <223> OTHER INFORMATION: The 'Xaa' at location 435 stands for Lys, Asn, Arg, Ser,
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     285   Tyr, Trp, Cys, or Phe.
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     288 <221> NAME/KEY: misc_feature
     289 <222> LOCATION: (436)..(436)
     290 <223> OTHER INFORMATION: The 'Xaa' at location 436 stands for Lys, Asn, Arg, Ser,
Thr, Ile
     291   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,
     292   Tyr, Trp, Cys, or Phe.
     294 <220> FEATURE:
     295 <221> NAME/KEY: misc_feature
     296 <222> LOCATION: (437)..(437)
     297 <223> OTHER INFORMATION: The 'Xaa' at location 437 stands for Lys, Asn, Arg, Ser,
Thr, Ile
     298   , Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon,

Please correct this error throughout Sequence 2

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299        ·Tyr, Trp, Cys, or Phe.
301 <220> FEATURE:
302 <221> NAME/KEY: misc_feature
303 <222> LOCATION: (438)..(438)
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 06/04/2002
PATENT APPLICATION: US/10/018,383   TIME: 14:56:07

Input Set : A:\X11600.ST25.txt
Output Set: N:\CRF3\06042002\J018383.raw

<u>Please Note:</u>
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:1; N Pos. 1333,1334,1335,1336,1337,1338,1339,1340,1341,1342,1343,1344
Seq#:1; N Pos. 1345,1346,1347,1348,1349,1350,1351,1352,1353,1354,1355,1356
Seq#:1; N Pos. 1357,1358,1359,1360,1361,1362,1363,1364,1365
Seq#:1; Xaa Pos. 421,422,423,424,425,426,427,428,429,430,431,432,433,434
Seq#:1; Xaa Pos. 435,436,437,438,439,440,441,442,443,444,445,446,447
Seq#:2; Xaa Pos. 421,422,423,424,425,426,427,428,429,430,431,432,433,434
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